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Crystal structure of enoyl-coenzyme A (CoA) hydratase at 2.5 angstroms resolution: a spiral fold defines the CoA-binding pocket.

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#	Paper	IF	Citations
155	The diverse world of coenzyme A binding proteins. 1996 , 6, 790-7		67
154	Paper Alert. <i>Structure</i> , 1996 , 4, 1531-1534	5.2	
153	Conformational heterogeneity in crystalline proteins: time-resolved fluorescence studies. 1997 , 278, 202-21		15
152	Glutamate-119 of the large alpha-subunit is the catalytic base in the hydration of 2-trans-enoyl-coenzyme A catalyzed by the multienzyme complex of fatty acid oxidation from <i>Escherichia coli</i> . <i>Biochemistry</i> , 1997 , 36, 11044-9	3.2	20
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